STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source:

Date Processed by STIC:

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,

2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 4.4.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
 U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street,
 Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

	
ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 10 6 8 5 5
ATTN: NEW RULES CASES:	PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE
1Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters , instead.
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5Variable Length	Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped
	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
8Skipped Sequences (NEW RULES)	Sequence(s) missing. If intentional , please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000
9Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
10Invalid 213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or Artificial Sequence
11 Use of <220>	Sequence(s) missing the <220> Teature and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
12PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13 Misuse of n/Xaa	"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



IFW16

RAW SEQUENCE LISTING

DATE: 03/19/2007

PATENT APPLICATION: US/10/618,852

TIME: 14:33:19

Input Set : N:\efs\03_19 07\10618852 efs\TVGN334ST25.txt

Output Set: N:\CRF4\03192007\J618852.raw

3 <110> APPLICANT: INVITROGEN CORPORATION 5 <120> TITLE OF INVENTION: SUBSCRIPTION BASED SYSTEMS, METHODS AND COMPONENTS FOR PROVIDING GENOMIC AND PROTEOMIC PRODUCTS AND SERVICES 6 8 <130> FILE REFERENCE: IVGN 334 10 <140> CURRENT APPLICATION NUMBER: 10/618,852 11 <141> CURRENT FILING DATE: 2003-07-15 13 <150> PRIOR APPLICATION NUMBER: 60396241 dres Not Comply 14 <151> PRIOR FILING DATE: 2002-07-17 16 <160> NUMBER OF SEQ ID NOS: 43 18 <170> SOFTWARE: PatentIn version 3.3 20 <210> SEO ID NO: 1 21 <211> LENGTH: 15 22 <212> TYPE: DNA 23 <213> ORGANISM: /Artificia 25 <220> FEATURE: 26 <223> OTHER INFORMATION: Wild-type att site 28 <400> SEQUENCE: 1 29 gcttttttat actaa 32 <210> SEQ ID NO: 2 33 <211> LENGTH: 21 34 <212> TYPE: DNA 35 <213> ORGANISM Artificial 37 <220> FEATURE: 38 <223> OTHER INFORMATION: Artificial sequence 40 <400> SEQUENCE: 2 41 caactttttt atacaaagtt g 44 <210> SEO ID NO: 3 45 <211> LENGTH: 25 46 <212> TYPE: DNA 47 <213> ORGANISM: Artificial 49 <220> FEATURE: 50 <223> OTHER INFORMATION: 52 <400> SEQUENCE: 3 53 agcctgcttt tttgtacaaa cttgt 25 56 <210> SEQ ID NO: 4 57 <211> LENGTH: 233 58 <212> TYPE: DNA 59 <213> ORGANISM: Artificial 61 <220> FEATURE: 62 <223 > OTHER INFORMATION 64 <400> SEQUENCE: 4 65 tacaggtcac taataccatc taagtagttg attcatagtg actggatatg ttgtgtttta 60

67 cagtattatg tagtctgttt tttatgcaaa atctaattta atatattgat atttatatca

120

DATE: 03/19/2007

TIME: 14:33:19

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                Output Set: N:\CRF4\03192007\J618852.raw
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71 tcaatttgtt gcaacgaaca ggtcactatc agtcaaaata aaatcattat ttq
74 <210> SEQ ID NO: 5
75 <211> LENGTH: 100
76 <212> TYPE: DNA
                                        same error
77 <213> ORGANISM: Artificial
79 <220> FEATURE:
80 <223> OTHER INFORMATION:
82 <400> SEQUENCE: 5
83 caaataatga ttttattttg actgatagtg acctgttcgt tgcaacaaat tgataagcaa
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85 tgcttttta taatgccaac tttgtacaaa aaagcaggct
                                                                         100
88 <210> SEQ ID NO: 6
89 <211> LENGTH: 125
90 <212> TYPE: DNA
                                       Same error
91 <213> ORGANISM: Artificial
93 <220> FEATURE:
94 <223> OTHER INFORMATION: /attr1
96 <400> SEQUENCE: 6
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97. acabattat acaaaaaago tgaacqaqaa acqtaaaaatq atataaatat caatatatta
99 aattagattt tgcataaaaa acagactaca taatactgta aaacacaaca tatccagtca
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101 ctatq
                                                                          125
104 <210> SEQ ID NO: 7
105 <211> LENGTH: 4
106 <212> TYPE: PRT
107 <213> ORGANISM: Artificial
109 <220> FEATURE:
110 <223> OTHER INFORMATION: Factor Xa cleavage site
112 <400> SEQUENCE: 7'
114 Ile Glu Gly Arg
115 1
118 <210> SEQ ID NO: 8
119 <211> LENGTH: 4
120 <212> TYPE: PRT
121 <213> ORGANISM: Artificial
123 <220> FEATURE:
124 <223> OTHER INFORMATION: Thrombin cleavage site
126 <400> SEQUENCE: 8
128 Leu Val Pro Arg
129 1
132 <210> SEQ ID NO: 9
133 <211> LENGTH: 27
134 <212> TYPE: DNA
135 <213> ORGANISM: Artificial
137 <220> FEATURE:
138 <223> OTHER INFORMATION!
140 <400> SEQUENCE: 9
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144 <210> SEQ ID NO: 10
145 <211> LENGTH: 27
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/618,852

RAW SEQUENCE LISTING DATE: 03/19/2007
PATENT APPLICATION: US/10/618,852 TIME: 14:33:19

Input Set : N:\efs\03_19_07\10618852_efs\IVGN334ST25.txt

Output Set: N:\CRF4\03192007\J618852.raw

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RAW SEQUENCE LISTING DATE: 03/19/2007
PATENT APPLICATION: US/10/618,852 TIME: 14:33:19

Input Set : N:\efs\03 19 07\10618852 efs\IVGN334ST25.txt

Output Set: N:\CRF4\03192007\J618852.raw

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291 <213> ORGANISM: Artificial

290 <212> TYPE: DNA

RAW SEQUENCE LISTING DATE: 03/19/2007 PATENT APPLICATION: US/10/618,852 TIME: 14:33:19

Input Set : N:\efs\03 19 07\10618852 efs\IVGN334ST25.txt

Output Set: N:\CRF4\03192007\J618852.raw

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 03/19/2007 PATENT APPLICATION: US/10/618,852 TIME: 14:33:20

Input Set : N:\efs\03_19_07\10618852_efs\IVGN334ST25.txt

Output Set: N:\CRF4\03192007\J618852.raw

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27 Seq#:28,29,30,31,32,33,34,35,36,37,38,39,40,41,42,43 VERIFICATION SUMMARY DATE: 03/19/2007 PATENT APPLICATION: US/10/618,852 TIME: 14:33:20

Input Set : N:\efs\03_19_07\10618852_efs\IVGN334ST25.txt
Output Set: N:\CRF4\03192007\J618852.raw